



FIG-1

FIG-2

FIG-3

Confirm or Enter Trimming Positions

Insertion Orientation

Positive

Negative

Starting or 5' Adapter/Vector Information

Sequence

Score

#

AGCGGCGCGCGGCGAGGT

68%

0

0

AGCGTGGTCCGCGCGCGAGGT

100%

0

0

Ending or 3' Adapter/Vector Information

Sequence

Score

#

ACCTCGGCGCGGACGCGCT

70%

0

0

ACCTGCGCGGGCGGCGCGCT

100%

0

0

TTACCGTGGTCCGCGCGCGAGGTACACAACACTGACAAACAAGGAAAGCGGAGAGAGAAAGACCTAGAGGTTGTAGATGGGAAATCAGGAATGATTTGAAGTAAAGATTTCCGGA
CTCATAAGAACACACATTTTAAATGTTAAACACAAAAACTACATGACTGAAGATAGAGAGAAATGCGATGGATTTTATACACATGGTGAAGAGAGAGAGGCGGTAGGTTTGCAAAACAAA
GTTAAGAAATAGGAAACTGAAATTTTTCATTGTACAGAAAATGTATCTCTTGGGAAAGGCGCTGTGACCTGCGCGCGCGCGCTCGAAATTCAGCACACTGCGCGCGCGCTTACTAGTGGATCC
CANCTCGGTACCAAGCTTGGAGTTATCATGGTCNTAANCCTGTTCCGTGTGAAATGTATCCNCCCATTCGCCCCACNTTCCAAACCGGAAACCTTAAATTTTAAACGNGGGTGCCN
AATGAATNACCGACCCCNNTATGCTTTGCCNCCCTGCCCTTCNCTCGGAACCNCTNCCCCTCTTNTAAACCGGCGCCCGGAAAGGTTGTTTGGCGCCCTCCCTGCCCTCCTCTNAT
CCTGCCCT

12

14

16

5' 4

3' 4

\\\\Mricd-fs1\sweeneyre\Sub Lib Test\Sequence\test\41x06.seq

Remove below base: 23

Length: 289

Confirm Fwd: TTACTAGTGGATCCGAGCTCGGTACCAAGCTT

0%

The current sequence was previously trimmed

Remove above base: 312

Threshold: 80

Confirm Rev: TTCGAACCATGGCTCGAGCC

0%

☒ Allow Inserts

☐ Positive Orientation

☒ Allow Deletes

☒ Use Threshold

Sequencing direction was not determined

Back

Apply

Skip

Abort All

Find Best